



PCT09

# ENTERED

## RAW SEQUENCE LISTING

DATE: 03/05/2002

PATENT APPLICATION: US/09/807,576

TIME: 13:39:13

Input Set : A:\USSN09807576.txt

Output Set: N:\CRF3\03052002\I807576.raw

3 &lt;110&gt; APPLICANT: ARIAD Gene Therapeutics, Inc.

5 &lt;120&gt; TITLE OF INVENTION: Materials and Methods Involving Conditional Aggregation

## Domains

7 &lt;130&gt; FILE REFERENCE: ARIAD 384A PCT/US

9 &lt;140&gt; CURRENT APPLICATION NUMBER: US 09/807,576

10 &lt;141&gt; CURRENT FILING DATE: 2001-04-13

12 &lt;160&gt; NUMBER OF SEQ ID NOS: 22

14 &lt;170&gt; SOFTWARE: PatentIn version 3.0

16 &lt;210&gt; SEQ ID NO: 1

17 &lt;211&gt; LENGTH: 7

18 &lt;212&gt; TYPE: PRT

19 &lt;213&gt; ORGANISM: Simian virus 40

21 &lt;220&gt; FEATURE:

22 &lt;221&gt; NAME/KEY: DOMAIN

23 &lt;222&gt; LOCATION: (1)..(7)

24 &lt;223&gt; OTHER INFORMATION: SV40 Large T nuclear localization domain

27 &lt;400&gt; SEQUENCE: 1

29 Pro Lys Lys Lys Arg Lys Val

30 1 5

32 &lt;210&gt; SEQ ID NO: 2

33 &lt;211&gt; LENGTH: 14

34 &lt;212&gt; TYPE: PRT

35 &lt;213&gt; ORGANISM: homo sapiens

37 &lt;220&gt; FEATURE:

38 &lt;221&gt; NAME/KEY: DOMAIN

39 &lt;222&gt; LOCATION: (1)..(14)

40 &lt;223&gt; OTHER INFORMATION: lipid targeting motif

43 &lt;400&gt; SEQUENCE: 2

45 Met Gly Ser Ser Lys Ser Lys Pro Lys Asp Pro Ser Gln Arg

46 1 5 10

48 &lt;210&gt; SEQ ID NO: 3

49 &lt;211&gt; LENGTH: 40

50 &lt;212&gt; TYPE: DNA

51 &lt;213&gt; ORGANISM: Artificial Sequence

53 &lt;220&gt; FEATURE:

54 &lt;221&gt; NAME/KEY: misc\_feature

55 &lt;222&gt; LOCATION: (1)..(40)

56 &lt;223&gt; OTHER INFORMATION: VR65 cloning oligo

59 &lt;400&gt; SEQUENCE: 3

60 tcccgccacct cttcggccag cgaattccag aagcgcgtat

40

63 &lt;210&gt; SEQ ID NO: 4

64 &lt;211&gt; LENGTH: 32

65 &lt;212&gt; TYPE: DNA

66 &lt;213&gt; ORGANISM: Artificial Sequence

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68 <220> FEATURE:
69 <221> NAME/KEY: misc_feature
70 <222> LOCATION: (1)..(32)
71 <223> OTHER INFORMATION: VR119 cloning oligo
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75 gactcactat aggacgcgtt cgagctcgcc cc 32
78 <210> SEQ ID NO: 5
79 <211> LENGTH: 31
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <221> NAME/KEY: misc_feature
85 <222> LOCATION: (1)..(31)
86 <223> OTHER INFORMATION: VR120 cloning oligo
89 <400> SEQUENCE: 5
90 catcattttg gcaaaggatt cactcctcag g 31
93 <210> SEQ ID NO: 6
94 <211> LENGTH: 27
95 <212> TYPE: DNA
96 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <221> NAME/KEY: misc_feature
100 <222> LOCATION: (1)..(27)
101 <223> OTHER INFORMATION: VR1 cloning oligo
104 <400> SEQUENCE: 6
105 gatggaaaaga aaatggattc ctcccgg 27
108 <210> SEQ ID NO: 7
109 <211> LENGTH: 24
110 <212> TYPE: DNA
111 <213> ORGANISM: Artificial Sequence
113 <220> FEATURE:
114 <221> NAME/KEY: misc_feature
115 <222> LOCATION: (1)..(24)
116 <223> OTHER INFORMATION: VR2 cloning oligo
119 <400> SEQUENCE: 7
120 tctagagtga gcaagggcga ggag 24
123 <210> SEQ ID NO: 8
124 <211> LENGTH: 37
125 <212> TYPE: DNA
126 <213> ORGANISM: Artificial Sequence
128 <220> FEATURE:
129 <221> NAME/KEY: misc_feature
130 <222> LOCATION: (1)..(37)
131 <223> OTHER INFORMATION: VR3 cloning oligo
134 <400> SEQUENCE: 8
135 ggatccttat taactagtct tgtacagctc gtccatg 37
138 <210> SEQ ID NO: 9
139 <211> LENGTH: 25
140 <212> TYPE: DNA

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141 <213> ORGANISM: Artificial Sequence
143 <220> FEATURE:
144 <221> NAME/KEY: misc_feature
145 <222> LOCATION: (1)..(25)
146 <223> OTHER INFORMATION: VR109 cloning oligo
149 <400> SEQUENCE: 9
150 aagcttacca ctcaggggtcc tgtgg                25
153 <210> SEQ ID NO: 10
154 <211> LENGTH: 19
155 <212> TYPE: DNA
156 <213> ORGANISM: Artificial Sequence
158 <220> FEATURE:
159 <221> NAME/KEY: misc_feature
160 <222> LOCATION: (1)..(19)
161 <223> OTHER INFORMATION: VR110 cloning oligo
164 <400> SEQUENCE: 10
165 gaattcgtgg caacttcca                19
168 <210> SEQ ID NO: 11
169 <211> LENGTH: 29
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171 <213> ORGANISM: Artificial Sequence
173 <220> FEATURE:
174 <221> NAME/KEY: misc_feature
175 <222> LOCATION: (1)..(29)
176 <223> OTHER INFORMATION: VR185 cloning oligo
179 <400> SEQUENCE: 11
180 cacaggaccc tgaattctaa gcttgtggc                29
183 <210> SEQ ID NO: 12
184 <211> LENGTH: 30
185 <212> TYPE: DNA
186 <213> ORGANISM: Artificial Sequence
188 <220> FEATURE:
189 <221> NAME/KEY: misc_feature
190 <222> LOCATION: (1)..(30)
191 <223> OTHER INFORMATION: VR186 cloning oligo
194 <400> SEQUENCE: 12
195 ataagggaat ggttctagag gcactgccct                30
198 <210> SEQ ID NO: 13
199 <211> LENGTH: 31
200 <212> TYPE: DNA
201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <221> NAME/KEY: misc_feature
205 <222> LOCATION: (1)..(31)
206 <223> OTHER INFORMATION: VR187 cloning oligo
209 <400> SEQUENCE: 13
210 atgccacccg ggactagtga agccacagct g                31
213 <210> SEQ ID NO: 14
214 <211> LENGTH: 48

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215 <212> TYPE: DNA
216 <213> ORGANISM: Artificial Sequence
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219 <221> NAME/KEY: misc_feature
220 <222> LOCATION: (1)..(48)
221 <223> OTHER INFORMATION: VR4 cloning oligo
224 <400> SEQUENCE: 14
225 actagtgccta gaaaccgtca gaagagattc ccaaccattc ccttaagc      48
228 <210> SEQ ID NO: 15
229 <211> LENGTH: 31
230 <212> TYPE: DNA
231 <213> ORGANISM: Artificial Sequence
233 <220> FEATURE:
234 <221> NAME/KEY: misc_feature
235 <222> LOCATION: (1)..(31)
236 <223> OTHER INFORMATION: VR5 cloning oligo
239 <400> SEQUENCE: 15
240 ggatcccgga ctagaagcca cagctgcoct c      31
243 <210> SEQ ID NO: 16
244 <211> LENGTH: 32
245 <212> TYPE: DNA
246 <213> ORGANISM: Artificial Sequence
248 <220> FEATURE:
249 <221> NAME/KEY: misc_feature
250 <222> LOCATION: (1)..(32)
251 <223> OTHER INFORMATION: VR 220 cloning oligo
254 <400> SEQUENCE: 16
255 cgaattcttc tgccatggcc ctgtggatgc gc      32
258 <210> SEQ ID NO: 17
259 <211> LENGTH: 31
260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <221> NAME/KEY: misc_feature
265 <222> LOCATION: (1)..(31)
266 <223> OTHER INFORMATION: VR221 cloning oligo
269 <400> SEQUENCE: 17
270 cggatccgca ggcgtgcgtct agttgcagta g      31
273 <210> SEQ ID NO: 18
274 <211> LENGTH: 52
275 <212> TYPE: DNA
276 <213> ORGANISM: Artificial Sequence
278 <220> FEATURE:
279 <221> NAME/KEY: misc_feature
280 <222> LOCATION: (1)..(52)
281 <223> OTHER INFORMATION: VR222 cloning oligo
284 <400> SEQUENCE: 18
285 cactagtgtc agaaaccgtc agaagagatt tgtgaaccaa cacctgtgcg gc      52
288 <210> SEQ ID NO: 19

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289 <211> LENGTH: 27
290 <212> TYPE: DNA
291 <213> ORGANISM: Artificial Sequence
293 <220> FEATURE:
294 <221> NAME/KEY: misc_feature
295 <222> LOCATION: (1)..(27)
296 <223> OTHER INFORMATION: VR223
299 <400> SEQUENCE: 19
300 cctgtgcggc tcagacctgg tggaagc 27
303 <210> SEQ ID NO: 20
304 <211> LENGTH: 35
305 <212> TYPE: DNA
306 <213> ORGANISM: Artificial Sequence
308 <220> FEATURE:
309 <221> NAME/KEY: misc_feature
310 <222> LOCATION: (1)..(35)
311 <223> OTHER INFORMATION: VR224 cloning oligo
314 <400> SEQUENCE: 20
315 cttctacaca cccaggacca agcgggaggc agagg 35
318 <210> SEQ ID NO: 21
319 <211> LENGTH: 29
320 <212> TYPE: DNA
321 <213> ORGANISM: Artificial Sequence
323 <220> FEATURE:
324 <221> NAME/KEY: misc_feature
325 <222> LOCATION: (1)..(29)
326 <223> OTHER INFORMATION: VR225 cloning oligo
329 <400> SEQUENCE: 21
330 ccctggaggg gtcccggcag aagcgtggc 29
333 <210> SEQ ID NO: 22
334 <211> LENGTH: 30
335 <212> TYPE: DNA
336 <213> ORGANISM: Artificial Sequence
338 <220> FEATURE:
339 <221> NAME/KEY: misc_feature
340 <222> LOCATION: (1)..(30)
341 <223> OTHER INFORMATION: oligo TC-W59V
344 <400> SEQUENCE: 22
345 gtgatccgag gcgtggaaga aggggttgcc 30

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VERIFICATION SUMMARY

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